

## CoNVex Name Clarification – June 2013

CoNVex (developed at the Wellcome Trust Sanger Institute) has been publicly available from our FTP repository for over 18 months and its name has been published in multiple conferences (including ASHG 2012, Genomic Disorder 2012, etc.). Another algorithm has been recently published (Amarasinge et al., 2013) with the same name in BMC Bioinformatics without being aware of our software, however its authors and the journal's editorial board have now changed their name to 'ADTEx' and published a revised manuscript to avoid the name conflict. References to the original and correction articles are available below.

We sincerely thank the authors of ADTEx and the BMC Bioinformatics' editorial board for their cooperation and assistance. We recommend the users of our software to confidently use the name 'CoNVex' while referring to our software.

### References

#### Original Article:

Amarasinghe K, Li J, Halgamuge S: CoNVEX: copy number variation estimation in exome sequencing data using HMM. BMC Bioinformatics 2013, 14(Suppl 2).

<http://www.biomedcentral.com/1471-2105/14/S2/S2>

#### Correction Article:

<http://www.biomedcentral.com/1471-2105/14/S2/S26>

Best wishes,  
Parthiban Vijayarangakannan.